

SEQUENCE LISTING

<110> Kodama, Tatsuhiko
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<120> Nonhuman animals for antibody production, and methods and systems for producing antibodies using such animals

<130> 14875-167US1

<150> PCT/JP2005/006298

<151> 2005-03-31

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<151> 2004-03-31

<160> 10

<170> PatentIn version 3.1

<210> 1

<211> 1539

<212> DNA

<213> Baculovirus

<220>

<221> CDS

<222> (1)..(1539)

<400> 1

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| 1 5 10 15 | |
| tct gcc ttt gcg gcg gag cac tgc aac gcg caa atg aag acg ggt ccg | 96 |
| Ser Ala Phe Ala Ala Glu His Cys Asn Ala Gln Met Lys Thr Gly Pro | |
| 20 25 30 | |
| tac aag att aaa aac ttg gac att acc ccg ccc aag gaa acg ctg caa | 144 |
| Tyr Lys Ile Lys Asn Leu Asp Ile Thr Pro Pro Lys Glu Thr Leu Gln | |
| 35 40 45 | |
| aag gac gtg gaa atc acc atc gtg gag acg gac tac aac gaa aac gtg | 192 |
| Lys Asp Val Glu Ile Thr Ile Val Glu Thr Asp Tyr Asn Glu Asn Val | |
| 50 55 60 | |
| att atc ggc tac aag ggg tac tac cag gcg tat gcg tac aac ggc ggc | 240 |
| Ile Ile Gly Tyr Lys Gly Tyr Tyr Gln Ala Tyr Ala Tyr Asn Gly Gly | |
| 65 70 75 80 | |
| tcg ctg gat ccc aac aca cgc gtc gaa gaa acc atg aaa acg ctg aat | 288 |
| Ser Leu Asp Pro Asn Thr Arg Val Glu Glu Thr Met Lys Thr Leu Asn | |
| 85 90 95 | |

| | |
|-----------------------------------------------------------------|-----|
| gtg ggc aaa gag gat ttg ctt atg tgg agc atc agg cag cag tgc gag | 336 |
| Val Gly Lys Glu Asp Leu Leu Met Trp Ser Ile Arg Gln Gln Cys Glu | |
| 100 105 110 | |
| gtg ggc gaa gag ctg atc gac cgt tgg ggc agt gac agc gac gac tgt | 384 |
| Val Gly Glu Glu Leu Ile Asp Arg Trp Gly Ser Asp Ser Asp Cys | |
| 115 120 125 | |
| ttt cgc gac aac gag ggc cgc ggc cag tgg gtc aaa ggc aaa gag ttg | 432 |
| Phe Arg Asp Asn Glu Gly Arg Gly Gln Trp Val Lys Gly Lys Glu Leu | |
| 130 135 140 | |
| gtg aag cgg cag aat aac aat cac ttt cgc cac cac acg tgc aac aaa | 480 |
| Val Lys Arg Gln Asn Asn Asn His Phe Ala His His Thr Cys Asn Lys | |
| 145 150 155 160 | |
| tcg tgg cga tgc ggc att tcc act tcg aaa atg tac agc agg ctc gag | 528 |
| Ser Trp Arg Cys Gly Ile Ser Thr Ser Lys Met Tyr Ser Arg Leu Glu | |
| 165 170 175 | |
| tgc cag gac gac acg gac gag tgc cag gta tac att ttg gac gct gag | 576 |
| Cys Gln Asp Asp Thr Asp Glu Cys Gln Val Tyr Ile Leu Asp Ala Glu | |
| 180 185 190 | |
| ggc aac ccc atc aac gtg acc gtg gac act gtg ctt cat cga gac ggc | 624 |
| Gly Asn Pro Ile Asn Val Thr Val Asp Thr Val Leu His Arg Asp Gly | |
| 195 200 205 | |
| gtg agt atg att ctc aaa caa aag tct acg ttc acc acg cgc caa ata | 672 |
| Val Ser Met Ile Leu Lys Gln Lys Ser Thr Phe Thr Thr Arg Gln Ile | |
| 210 215 220 | |
| aaa gct cgc tgt ctg ctc att aaa gat gac aaa aat aac ccc gag tcg | 720 |
| Lys Ala Ala Cys Leu Leu Ile Lys Asp Asp Lys Asn Asn Pro Glu Ser | |
| 225 230 235 240 | |
| gtg aca cgc gaa cac tgt ttg att gac aat gat ata tat gat ctt tct | 768 |
| Val Thr Arg Glu His Cys Leu Ile Asp Asn Asp Ile Tyr Asp Leu Ser | |
| 245 250 255 | |
| aaa aac acg tgg aac tgc aag ttt aac aga tgc att aaa cgc aaa gtc | 816 |
| Lys Asn Thr Trp Asn Cys Lys Phe Asn Arg Cys Ile Lys Arg Lys Val | |
| 260 265 270 | |
| gag cac cga gtc aag aag cgg ccg ccc act tgg cgc cac aac gtt aga | 864 |
| Glu His Arg Val Lys Lys Arg Pro Pro Thr Trp Arg His Asn Val Arg | |
| 275 280 285 | |
| gcc aag tac aca gag gga gac act gcc acc aaa ggc gac ctg atg cat | 912 |
| Ala Lys Tyr Thr Glu Gly Asp Thr Ala Thr Lys Gly Asp Leu Met His | |
| 290 295 300 | |
| att caa gag gag ctg atg tac gaa aac gat ttg ctg aaa atg aac att | 960 |
| Ile Gln Glu Glu Leu Met Tyr Glu Asn Asp Leu Leu Lys Met Asn Ile | |
| 305 310 315 320 | |

| | |
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| gag ctg atg cat gcg cac atc aac aag cta aac aat atg ctg cac gac Glu Leu Met His Ala His Ile Asn Lys Leu Asn Asn Met Leu His Asp 325 330 335 | 1008 |
| ctg ata gtc tcc gtg gcc aag gtg gac gag cgt ttg att ggc aat ctc Leu Ile Val Ser Val Ala Lys Val Asp Glu Arg Leu Ile Gly Asn Leu 340 345 350 | 1056 |
| atg aac aac tct gtt tct tca aca ttt ttg tcy gac gac acg ttt ttg Met Asn Asn Ser Val Ser Ser Thr Phe Leu Ser Asp Asp Thr Phe Leu 355 360 365 | 1104 |
| ctg atg ccg tgc acc aat ccg ccg gca cac acc agt aat tgc tac aac Leu Met Pro Cys Thr Asn Pro Pro Ala His Thr Ser Asn Cys Tyr Asn 370 375 380 | 1152 |
| aac agc atc tac aaa gaa ggg cgt tgg gtg gcc aac acg gac tcy tcy Asn Ser Ile Tyr Lys Glu Gly Arg Trp Val Ala Asn Thr Asp Ser Ser 385 390 395 400 | 1200 |
| caa tgc ata gat ttt agc aac tac aag gaa cta gca att gac gac gac Gln Cys Ile Asp Phe Ser Asn Tyr Lys Glu Leu Ala Ile Asp Asp Asp 405 410 415 | 1248 |
| gtc gag ttt tgg atc ccg acc atc ggc aac acg acc tat cac gac agt Val Glu Phe Trp Ile Pro Thr Ile Gly Asn Thr Thr Tyr His Asp Ser 420 425 430 | 1296 |
| tgg aaa gat gcc agc ggc tgg tcy ttt att gcc caa caa aaa agc aac Trp Lys Asp Ala Ser Gly Trp Ser Phe Ile Ala Gln Gln Lys Ser Asn 435 440 445 | 1344 |
| ctc ata acc acc atg gag aac acc aag ttt ggc ggc gtc ggc acc agt Leu Ile Thr Thr Met Glu Asn Thr Lys Phe Gly Gly Val Gly Thr Ser 450 455 460 | 1392 |
| ctg agc gac atc act tcc atg gct gaa ggc gaa ttg gcc gct aaa ttg Leu Ser Asp Ile Thr Ser Met Ala Glu Gly Glu Leu Ala Ala Lys Leu 465 470 475 480 | 1440 |
| act tcy ttc atg ttt ggt cat gta gtt aac ttt gta att ata tta att Thr Ser Phe Met Phe Gly His Val Val Asn Phe Val Ile Ile Leu Ile 485 490 495 | 1488 |
| gtg att tta ttt ttg tac tgt atg att aga aac cgt aat aga caa tat Val Ile Leu Phe Leu Tyr Cys Met Ile Arg Asn Arg Asn Arg Gln Tyr 500 505 510 | 1536 |
| taa | 1539 |

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 <213> Baculovirus
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 Ser Ala Phe Ala Ala Glu His Cys Asn Ala Gln Met Lys Thr Gly Pro
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 Tyr Lys Ile Lys Asn Leu Asp Ile Thr Pro Pro Lys Glu Thr Leu Gln
 35 40 45
 Lys Asp Val Glu Ile Thr Ile Val Glu Thr Asp Tyr Asn Glu Asn Val
 50 55 60
 Ile Ile Gly Tyr Lys Gly Tyr Tyr Gln Ala Tyr Ala Tyr Asn Gly Gly
 65 70 75 80
 Ser Leu Asp Pro Asn Thr Arg Val Glu Glu Thr Met Lys Thr Leu Asn
 85 90 95
 Val Gly Lys Glu Asp Leu Leu Met Trp Ser Ile Arg Gln Gln Cys Glu
 100 105 110
 Val Gly Glu Glu Leu Ile Asp Arg Trp Gly Ser Asp Ser Asp Asp Cys
 115 120 125
 Phe Arg Asp Asn Glu Gly Arg Gly Gln Trp Val Lys Gly Lys Glu Leu
 130 135 140
 Val Lys Arg Gln Asn Asn Asn His Phe Ala His His Thr Cys Asn Lys
 145 150 155 160
 Ser Trp Arg Cys Gly Ile Ser Thr Ser Lys Met Tyr Ser Arg Leu Glu
 165 170 175
 Cys Gln Asp Asp Thr Asp Glu Cys Gln Val Tyr Ile Leu Asp Ala Glu
 180 185 190
 Gly Asn Pro Ile Asn Val Thr Val Asp Thr Val Leu His Arg Asp Gly
 195 200 205
 Val Ser Met Ile Leu Lys Gln Lys Ser Thr Phe Thr Thr Arg Gln Ile
 210 215 220
 Lys Ala Ala Cys Leu Leu Ile Lys Asp Asp Lys Asn Asn Pro Glu Ser
 225 230 235 240
 Val Thr Arg Glu His Cys Leu Ile Asp Asn Asp Ile Tyr Asp Leu Ser
 245 250 255
 Lys Asn Thr Trp Asn Cys Lys Phe Asn Arg Cys Ile Lys Arg Lys Val
 260 265 270
 Glu His Arg Val Lys Lys Arg Pro Pro Thr Trp Arg His Asn Val Arg
 275 280 285
 Ala Lys Tyr Thr Glu Gly Asp Thr Ala Thr Lys Gly Asp Leu Met His
 290 295 300

Ile Gln Glu Glu Leu Met Tyr Glu Asn Asp Leu Leu Lys Met Asn Ile
305 310 315 320

Glu Leu Met His Ala His Ile Asn Lys Leu Asn Asn Met Leu His Asp
325 330 335

Leu Ile Val Ser Val Ala Lys Val Asp Glu Arg Leu Ile Gly Asn Leu
340 345 350

Met Asn Asn Ser Val Ser Ser Thr Phe Leu Ser Asp Asp Thr Phe Leu
355 360 365

Leu Met Pro Cys Thr Asn Pro Pro Ala His Thr Ser Asn Cys Tyr Asn
370 375 380

Asn Ser Ile Tyr Lys Glu Gly Arg Trp Val Ala Asn Thr Asp Ser Ser
385 390 395 400

Gln Cys Ile Asp Phe Ser Asn Tyr Lys Glu Leu Ala Ile Asp Asp Asp
405 410 415

Val Glu Phe Trp Ile Pro Thr Ile Gly Asn Thr Thr Tyr His Asp Ser
420 425 430

Trp Lys Asp Ala Ser Gly Trp Ser Phe Ile Ala Gln Gln Lys Ser Asn
435 440 445

Leu Ile Thr Thr Met Glu Asn Thr Lys Phe Gly Gly Val Gly Thr Ser
450 455 460

Leu Ser Asp Ile Thr Ser Met Ala Glu Gly Glu Leu Ala Ala Lys Leu
465 470 475 480

Thr Ser Phe Met Phe Gly His Val Val Asn Phe Val Ile Ile Leu Ile
485 490 495

Val Ile Leu Phe Leu Tyr Cys Met Ile Arg Asn Arg Asn Arg Gln Tyr
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<220>
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<222> (1)..(1464)

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tct gcc ttt gcg gcg gag cac tgc aac gcg caa atg aag acg ggt ccg 96
Ser Ala Phe Ala Ala Glu His Cys Asn Ala Gln Met Lys Thr Gly Pro
20 25 30

| | |
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| tac aag att aaa aac ttg gac att acc ccg ccc aag gaa acg ctg caa Tyr Lys Ile Lys Asn Leu Asp Ile Thr Pro Pro Lys Glu Thr Leu Gln 35 40 45 | 144 |
| aag gac gtg gaa atc acc atc gtg gag acg gac tac aac gaa aac gtg Lys Asp Val Glu Ile Thr Ile Val Glu Thr Asp Tyr Asn Glu Asn Val 50 55 60 | 192 |
| att atc ggc tac aag ggg tac tac cag gcg tat gcg tac aac ggc ggc Ile Ile Gly Tyr Lys Gly Tyr Tyr Gln Ala Tyr Ala Tyr Asn Gly Gly 65 70 75 80 | 240 |
| tcg ctg gat ccc aac aca cgc gtc gaa gaa acc atg aaa acg ctg aat Ser Leu Asp Pro Asn Thr Arg Val Glu Glu Thr Met Lys Thr Leu Asn 85 90 95 | 288 |
| gtg ggc aaa gag gat ttg ctt atg tgg agc atc agg cag cag tgc gag Val Gly Lys Glu Asp Leu Leu Met Trp Ser Ile Arg Gln Gln Cys Glu 100 105 110 | 336 |
| gtg ggc gaa gag ctg atc gac cgt tgg ggc agt gac agc gac gac tgt Val Gly Glu Glu Leu Ile Asp Arg Trp Gly Ser Asp Ser Asp Asp Cys 115 120 125 | 384 |
| ttt cgc gac aac gag ggc cgc ggc cag tgg gtc aaa ggc aaa gag ttg Phe Arg Asp Asn Glu Gly Arg Gly Gln Trp Val Lys Gly Lys Glu Leu 130 135 140 | 432 |
| gtg aag cgg cag aat aac aat cac ttt cgc cac cac acg tgc aac aaa Val Lys Arg Gln Asn Asn Asn His Phe Ala His His Thr Cys Asn Lys 145 150 155 160 | 480 |
| tcg tgg cga tgc ggc att tcc act tcg aaa atg tac agc agg ctc gag Ser Trp Arg Cys Gly Ile Ser Thr Ser Lys Met Tyr Ser Arg Leu Glu 165 170 175 | 528 |
| tgc cag gac gac acg gac gag tgc cag gta tac att ttg gac gct gag Cys Gln Asp Asp Thr Asp Glu Cys Gln Val Tyr Ile Leu Asp Ala Glu 180 185 190 | 576 |
| ggc aac ccc atc aac gtg acc gtg gac act gtg ctt cat cga gac ggc Gly Asn Pro Ile Asn Val Thr Val Asp Thr Val Leu His Arg Asp Gly 195 200 205 | 624 |
| gtg agt atg att ctc aaa caa aag tct acg ttc acc acg cgc caa ata Val Ser Met Ile Leu Lys Gln Lys Ser Thr Phe Thr Thr Arg Gln Ile 210 215 220 | 672 |
| aaa gct gcg tgt ctg ctc att aaa gat gac aaa aat aac ccc gag tcg Lys Ala Ala Cys Leu Leu Ile Lys Asp Lys Asn Asn Pro Glu Ser 225 230 235 240 | 720 |
| gtg aca cgc gaa cac tgt ttg att gac aat gat ata tat gat ctt tct Val Thr Arg Glu His Cys Leu Ile Asp Asn Asp Ile Tyr Asp Leu Ser 245 250 255 | 768 |

| | |
|-------------------------------------------------------------------------------------------------------------------------------------------------------|------|
| aaa aac acg tgg aac tgc aag ttt aac aga tgc att aaa cgc aaa gtc Lys Asn Thr Trp Asn Cys Lys Phe Asn Arg Cys Ile Lys Arg Lys Val 260 265 270 | 816 |
| gag cac cga gtc aag aag cgg ccg ccc act tgg cgc cac aac gtt aga Glu His Arg Val Lys Lys Arg Pro Pro Thr Trp Arg His Asn Val Arg 275 280 285 | 864 |
| gcc aag tac aca gag gga gac act gcc acc aaa ggc gac ctg atg cat Ala Lys Tyr Thr Glu Gly Asp Thr Ala Thr Lys Gly Asp Leu Met His 290 295 300 | 912 |
| att caa gag gag ctg atg tac gaa aac gat ttg ctg aaa atg aac att Ile Gln Glu Glu Leu Met Tyr Glu Asn Asp Leu Lys Leu Met Asn Ile 305 310 315 320 | 960 |
| gag ctg atg cat gcg cac atc aac aag cta aac aat atg ctg cac gac Glu Leu Met His Ala His Ile Asn Lys Leu Asn Asn Met Leu His Asp 325 330 335 | 1008 |
| ctg ata gtc tcc gtg gcc aag gtg gac gag cgt ttg att ggc aat ctc Leu Ile Val Ser Val Ala Lys Val Asp Glu Arg Leu Ile Gly Asn Leu 340 345 350 | 1056 |
| atg aac aac tct gtt tct tca aca ttt ttg tgc gac gac acg ttt ttg Met Asn Asn Ser Val Ser Ser Thr Phe Leu Ser Asp Asp Thr Phe Leu 355 360 365 | 1104 |
| ctg atg ccg tgc acc aat ccg ccg gca cac acc agt aat tgc tac aac Leu Met Pro Cys Thr Asn Pro Pro Ala His Thr Ser Asn Cys Tyr Asn 370 375 380 | 1152 |
| aac agc atc tac aaa gaa ggg cgt tgg gtg gcc aac acg gac tcg tcg Asn Ser Ile Tyr Lys Glu Gly Arg Trp Val Ala Asn Thr Asp Ser Ser 385 390 395 400 | 1200 |
| caa tgc ata gat ttt agc aac tac aag gaa cta gca att gac gac gac Gln Cys Ile Asp Phe Ser Asn Tyr Lys Glu Leu Ala Ile Asp Asp Asp 405 410 415 | 1248 |
| gtc gag ttt tgg atc ccg acc atc ggc aac acg acc tat cac gac agt Val Glu Phe Trp Ile Pro Thr Ile Gly Asn Thr Thr Tyr His Asp Ser 420 425 430 | 1296 |
| tgg aaa gat gcc agc ggc tgg tgc ttt att gcc caa caa aaa agc aac Trp Lys Asp Ala Ser Gly Trp Phe Ile Ala Gln Lys Ser Asn 435 440 445 | 1344 |
| ctc ata acc acc atg gag aac acc aag ttt ggc ggc gtc ggc acc agt Leu Ile Thr Thr Met Glu Asn Thr Lys Phe Gly Gly Val Gly Thr Ser 450 455 460 | 1392 |
| ctg agc gac atc act tcc atg gct gaa ggc gaa ttg gcc gct aaa ttg Leu Ser Asp Ile Thr Ser Met Ala Glu Gly Glu Leu Ala Ala Lys Leu 465 470 475 480 | 1440 |
| act tcg ttc atg ttt ggt cat gta | 1464 |

Thr Ser Phe Met Phe Gly His Val
485

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<213> Baculovirus

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Tyr Lys Ile Lys Asn Leu Asp Ile Thr Pro Pro Lys Glu Thr Leu Gln
35 40 45
Lys Asp Val Glu Ile Thr Ile Val Glu Thr Asp Tyr Asn Glu Asn Val
50 55 60
Ile Ile Gly Tyr Lys Gly Tyr Tyr Gln Ala Tyr Ala Tyr Asn Gly Gly
65 70 75 80
Ser Leu Asp Pro Asn Thr Arg Val Glu Glu Thr Met Lys Thr Leu Asn
85 90 95
Val Gly Lys Glu Asp Leu Leu Met Trp Ser Ile Arg Gln Gln Cys Glu
100 105 110
Val Gly Glu Glu Leu Ile Asp Arg Trp Gly Ser Asp Ser Asp Asp Cys
115 120 125
Phe Arg Asp Asn Glu Gly Arg Gly Gln Trp Val Lys Gly Lys Glu Leu
130 135 140
Val Lys Arg Gln Asn Asn Asn His Phe Ala His His Thr Cys Asn Lys
145 150 155 160
Ser Trp Arg Cys Gly Ile Ser Thr Ser Lys Met Tyr Ser Arg Leu Glu
165 170 175
Cys Gln Asp Asp Thr Asp Glu Cys Gln Val Tyr Ile Leu Asp Ala Glu
180 185 190
Gly Asn Pro Ile Asn Val Thr Val Asp Thr Val Leu His Arg Asp Gly
195 200 205
Val Ser Met Ile Leu Lys Gln Lys Ser Thr Phe Thr Thr Arg Gln Ile
210 215 220
Lys Ala Ala Cys Leu Leu Ile Lys Asp Asp Lys Asn Asn Pro Glu Ser
225 230 235 240
Val Thr Arg Glu His Cys Leu Ile Asp Asn Asp Ile Tyr Asp Leu Ser
245 250 255

Lys Asn Thr Trp Asn Cys Lys Phe Asn Arg Cys Ile Lys Arg Lys Val
 260 265 270
 Glu His Arg Val Lys Lys Arg Pro Pro Thr Trp Arg His Asn Val Arg
 275 280 285
 Ala Lys Tyr Thr Glu Gly Asp Thr Ala Thr Lys Gly Asp Leu Met His
 290 295 300
 Ile Gln Glu Glu Leu Met Tyr Glu Asn Asp Leu Lys Met Asn Ile
 305 310 315 320
 Glu Leu Met His Ala His Ile Asn Lys Leu Asn Asn Met Leu His Asp
 325 330 335
 Leu Ile Val Ser Val Ala Lys Val Asp Glu Arg Leu Ile Gly Asn Leu
 340 345 350
 Met Asn Asn Ser Val Ser Ser Thr Phe Leu Ser Asp Asp Thr Phe Leu
 355 360 365
 Leu Met Pro Cys Thr Asn Pro Pro Ala His Thr Ser Asn Cys Tyr Asn
 370 375 380
 Asn Ser Ile Tyr Lys Glu Gly Arg Trp Val Ala Asn Thr Asp Ser Ser
 385 390 395 400
 Gln Cys Ile Asp Phe Ser Asn Tyr Lys Glu Leu Ala Ile Asp Asp Asp
 405 410 415
 Val Glu Phe Trp Ile Pro Thr Ile Gly Asn Thr Thr Tyr His Asp Ser
 420 425 430
 Trp Lys Asp Ala Ser Gly Trp Ser Phe Ile Ala Gln Gln Lys Ser Asn
 435 440 445
 Leu Ile Thr Thr Met Glu Asn Thr Lys Phe Gly Gly Val Gly Thr Ser
 450 455 460
 Leu Ser Asp Ile Thr Ser Met Ala Glu Gly Glu Leu Ala Ala Lys Leu
 465 470 475 480
 Thr Ser Phe Met Phe Gly His Val
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33

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 <213> Artificial

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17

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<220>
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<400> 8
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24

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 <212> DNA
 <213> Baculovirus

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 <222> (2)..(1474)

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 Asn Ser Thr Met Val Ser Ala Ile Val Leu Tyr Val Leu Leu Ala Ala
 1 5 10 15

49

gcg gcg cat tct gcc ttt gcg gcg gag cac tgc aac gcg caa atg aag
 Ala Ala His Ser Ala Phe Ala Ala Glu His Cys Asn Ala Gln Met Lys
 20 25 30

97

| | |
|-----------------------------------------------------------------|-----|
| acg ggt ccg tac aag att aaa aac ttg gac att acc ccg ccc aag gaa | 145 |
| Thr Gly Pro Tyr Lys Ile Lys Asn Leu Asp Ile Thr Pro Pro Lys Glu | |
| 35 40 45 | |
| acg ctg caa aag gac gtg gaa atc acc atc gtg gag acg gac tac aac | 193 |
| Thr Leu Gln Lys Asp Val Glu Ile Thr Ile Val Glu Thr Asp Tyr Asn | |
| 50 55 60 | |
| gaa aac gtg att atc ggc tac aag ggg tac tac cag gcg tat gcg tac | 241 |
| Glu Asn Val Ile Ile Gly Tyr Lys Gly Tyr Tyr Gln Ala Tyr Ala Tyr | |
| 65 70 75 80 | |
| aac ggc ggc tcg ctg gat ccc aac aca cgc gtc gaa gaa acc atg aaa | 289 |
| Asn Gly Gly Ser Leu Asp Pro Asn Thr Arg Val Glu Glu Thr Met Lys | |
| 85 90 95 | |
| acg ctg aat gtg ggc aaa gag gat ttg ctt atg tgg agc atc agg cag | 337 |
| Thr Leu Asn Val Gly Lys Glu Asp Leu Leu Met Trp Ser Ile Arg Gln | |
| 100 105 110 | |
| cag tgc gag gtg ggc gaa gag ctg atc gac cgt tgg ggc agt gac agc | 385 |
| Gln Cys Glu Val Gly Glu Glu Leu Ile Asp Arg Trp Gly Ser Asp Ser | |
| 115 120 125 | |
| gac gac tgt ttt cgc gac aac gag ggc cgc ggc cag tgg gtc aaa ggc | 433 |
| Asp Asp Cys Phe Arg Asp Asn Glu Gly Arg Gly Gln Trp Val Lys Gly | |
| 130 135 140 | |
| aaa gag ttg gtg aag cgg cag aat aac aat cac ttt gcg cac cac acg | 481 |
| Lys Glu Leu Val Lys Arg Gln Asn Asn Asn His Phe Ala His His Thr | |
| 145 150 155 160 | |
| tgc aac aaa tcg tgg cga tgc ggc att tcc act tcg aaa atg tac agc | 529 |
| Cys Asn Lys Ser Trp Arg Cys Gly Ile Ser Thr Ser Lys Met Tyr Ser | |
| 165 170 175 | |
| agg ctc gag tgc cag gac gac acg gac gag tgc cag gta tac att ttg | 577 |
| Arg Leu Glu Cys Gln Asp Asp Thr Asp Glu Cys Gln Val Tyr Ile Leu | |
| 180 185 190 | |
| gac gct gag ggc aac ccc atc aac gtg acc gtg gac act gtg ctt cat | 625 |
| Asp Ala Glu Gly Asn Pro Ile Asn Val Thr Val Asp Thr Val Leu His | |
| 195 200 205 | |
| cga gac ggc gtg agt atg att ctc aaa caa aag tct acg ttc acc acg | 673 |
| Arg Asp Gly Val Ser Met Ile Leu Lys Gln Lys Ser Thr Phe Thr Thr | |
| 210 215 220 | |
| cgc caa ata aaa gct gcg tgt ctg ctc att aaa gat gac aaa aat aac | 721 |
| Arg Gln Ile Lys Ala Ala Cys Leu Leu Ile Lys Asp Asp Lys Asn Asn | |
| 225 230 235 240 | |
| ccc gag tcg gtg aca cgc gaa cac tgt ttg att gac aat gat ata tat | 769 |
| Pro Glu Ser Val Thr Arg Glu His Cys Leu Ile Asp Asn Asp Ile Tyr | |
| 245 250 255 | |

| | |
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| gat ctt tct aaa aac acg tgg aac tgc aag ttt aac aga tgc att aaa | 817 |
| Asp Leu Ser Lys Asn Thr Trp Asn Cys Lys Phe Asn Arg Cys Ile Lys | |
| 260 265 270 | |
| cgc aaa gtc gag cac cga gtc aag aag cgg ccg ccc act tgg cgc cac | 865 |
| Arg Lys Val Glu His Arg Val Lys Lys Arg Pro Pro Thr Trp Arg His | |
| 275 280 285 | |
| aac gtt aga gcc aag tac aca gag gga gac act gcc acc aaa ggc gac | 913 |
| Asn Val Arg Ala Lys Tyr Thr Glu Gly Asp Thr Ala Thr Lys Gly Asp | |
| 290 295 300 | |
| ctg atg cat att caa gag gag ctg atg tac gaa aac gat ttg ctg aaa | 961 |
| Leu Met His Ile Gln Glu Glu Leu Met Tyr Glu Asn Asp Leu Leu Lys | |
| 305 310 315 320 | |
| atg aac att gag ctg atg cat gcg cac atc aac aag cta aac aat atg | 1009 |
| Met Asn Ile Glu Leu Met His Ala His Ile Asn Lys Leu Asn Asn Met | |
| 325 330 335 | |
| ctg cac gac ctg ata gtc tcc gtg gcc aag gtg gac gag cgt ttg att | 1057 |
| Leu His Asp Leu Ile Val Ser Val Ala Lys Val Asp Glu Arg Leu Ile | |
| 340 345 350 | |
| ggc aat ctg atg aac aac tct gtt tct tca aca ttt ttg tcg gac gac | 1105 |
| Gly Asn Leu Met Asn Asn Ser Val Ser Ser Thr Phe Leu Ser Asp Asp | |
| 355 360 365 | |
| acg ttt ttg ctg atg ccg tgc acc aat ccg ccg gca cac acc agt aat | 1153 |
| Thr Phe Leu Leu Met Pro Cys Thr Asn Pro Pro Ala His Thr Ser Asn | |
| 370 375 380 | |
| tgc tac aac aac agc atc tac aaa gaa ggg cgt tgg gtg gcc aac acg | 1201 |
| Cys Tyr Asn Asn Ser Ile Tyr Lys Glu Gly Arg Trp Val Ala Asn Thr | |
| 385 390 395 400 | |
| gac tcg tcg caa tgc ata gat ttt agc aac tac aag gaa cta gca att | 1249 |
| Asp Ser Ser Gln Cys Ile Asp Phe Ser Asn Tyr Lys Glu Leu Ala Ile | |
| 405 410 415 | |
| gac gac gac gtc gag ttt tgg atc ccg acc atc ggc aac acg acc tat | 1297 |
| Asp Asp Asp Val Glu Phe Trp Ile Pro Thr Ile Gly Asn Thr Thr Tyr | |
| 420 425 430 | |
| cac gac agt tgg aaa gat gcc agc ggc tgg tcg ttt att gcc caa caa | 1345 |
| His Asp Ser Trp Lys Asp Ala Ser Gly Trp Ser Phe Ile Ala Gln Gln | |
| 435 440 445 | |
| aaa agc aac ctg ata acc acc atg gag aac acc aag ttt ggc ggc gtc | 1393 |
| Lys Ser Asn Leu Ile Thr Thr Met Glu Asn Thr Lys Phe Gly Gly Val | |
| 450 455 460 | |
| ggc acc agt ctg agc gac atc act tcc atg gct gaa ggc gaa ttg gcc | 1441 |
| Gly Thr Ser Leu Ser Asp Ile Thr Ser Met Ala Glu Gly Glu Leu Ala | |
| 465 470 475 480 | |
| gct aaa ttg act tcg ttc atg ttt ggt cat gta taatgagaat tc | 1486 |

Ala Lys Leu Thr Ser Phe Met Phe Gly His Val
 485 490

<210> 10
 <211> 491
 <212> PRT
 <213> Baculovirus

<400> 10

Asn Ser Thr Met Val Ser Ala Ile Val Leu Tyr Val Leu Leu Ala Ala
 1 5 10 15

Ala Ala His Ser Ala Phe Ala Ala Glu His Cys Asn Ala Gln Met Lys
 20 25 30

Thr Gly Pro Tyr Lys Ile Lys Asn Leu Asp Ile Thr Pro Pro Lys Glu
 35 40 45

Thr Leu Gln Lys Asp Val Glu Ile Thr Ile Val Glu Thr Asp Tyr Asn
 50 55 60

Glu Asn Val Ile Ile Gly Tyr Lys Gly Tyr Tyr Gln Ala Tyr Ala Tyr
 65 70 75 80

Asn Gly Gly Ser Leu Asp Pro Asn Thr Arg Val Glu Glu Thr Met Lys
 85 90 95

Thr Leu Asn Val Gly Lys Glu Asp Leu Leu Met Trp Ser Ile Arg Gln
 100 105 110

Gln Cys Glu Val Gly Glu Glu Leu Ile Asp Arg Trp Gly Ser Asp Ser
 115 120 125

Asp Asp Cys Phe Arg Asp Asn Glu Gly Arg Gly Gln Trp Val Lys Gly
 130 135 140

Lys Glu Leu Val Lys Arg Gln Asn Asn Asn His Phe Ala His His Thr
 145 150 155 160

Cys Asn Lys Ser Trp Arg Cys Gly Ile Ser Thr Ser Lys Met Tyr Ser
 165 170 175

Arg Leu Glu Cys Gln Asp Asp Thr Asp Glu Cys Gln Val Tyr Ile Leu
 180 185 190

Asp Ala Glu Gly Asn Pro Ile Asn Val Thr Val Asp Thr Val Leu His
195 200 205

Arg Asp Gly Val Ser Met Ile Leu Lys Gln Lys Ser Thr Phe Thr Thr
210 215 220

Arg Gln Ile Lys Ala Ala Cys Leu Leu Ile Lys Asp Asp Lys Asn Asn
225 230 235 240

Pro Glu Ser Val Thr Arg Glu His Cys Leu Ile Asp Asn Asp Ile Tyr
245 250 255

Asp Leu Ser Lys Asn Thr Trp Asn Cys Lys Phe Asn Arg Cys Ile Lys
260 265 270

Arg Lys Val Glu His Arg Val Lys Lys Arg Pro Pro Thr Trp Arg His
275 280 285

Asn Val Arg Ala Lys Tyr Thr Glu Gly Asp Thr Ala Thr Lys Gly Asp
290 295 300

Leu Met His Ile Gln Glu Glu Leu Met Tyr Glu Asn Asp Leu Leu Lys
305 310 315 320

Met Asn Ile Glu Leu Met His Ala His Ile Asn Lys Leu Asn Asn Met
325 330 335

Leu His Asp Leu Ile Val Ser Val Ala Lys Val Asp Glu Arg Leu Ile
340 345 350

Gly Asn Leu Met Asn Asn Ser Val Ser Ser Thr Phe Leu Ser Asp Asp
355 360 365

Thr Phe Leu Leu Met Pro Cys Thr Asn Pro Pro Ala His Thr Ser Asn
370 375 380

Cys Tyr Asn Asn Ser Ile Tyr Lys Glu Gly Arg Trp Val Ala Asn Thr
385 390 395 400

Asp Ser Ser Gln Cys Ile Asp Phe Ser Asn Tyr Lys Glu Leu Ala Ile
405 410 415

Asp Asp Asp Val Glu Phe Trp Ile Pro Thr Ile Gly Asn Thr Thr Tyr
 420 425 430

His Asp Ser Trp Lys Asp Ala Ser Gly Trp Ser Phe Ile Ala Gln Gln
 435 440 445

Lys Ser Asn Leu Ile Thr Thr Met Glu Asn Thr Lys Phe Gly Gly Val
 450 455 460

Gly Thr Ser Leu Ser Asp Ile Thr Ser Met Ala Glu Gly Glu Leu Ala
 465 470 475 480

Ala Lys Leu Thr Ser Phe Met Phe Gly His Val
 485 490